•	CRF Errors Corrected by the STIC Systems Branch  CRF Processing Date: 6/27/2002
	CRF Ert S Corrected by the STIC Systems Branch  CRF Processing Date: 6/21/2002
riai i	Edited by:
	Changed a file from non-ASCII to ASCIIENTED Verified by: Verified by: (STIC staff)
	Changed the margins in cases where the sequence text was "wrapped" down to the next line.
	Edited a format error in the Current Application Data section, specifically:
	Edited the Current Application Data section with the actual current number. The number inputted by the applicant was  the prior application data; or other
	Added the mandatory heading and subheadings for "Current Application Data".
	Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
	Changed the spelling of a mandatory field (the headings or subheadings), specifically:
; <del>-</del>	Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
	Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
	Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
	Inserted colons after headings/subheadings. Headings edited included:
	Deleted extra, invalid, headings used by an applicant, specifically:
	Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as
	Inserted mandatory headings, specifically:
	Corrected an obvious error in the response, specifically:
	Edited identifiers where upper case is used but lower case is required, or vice versa.
	Corrected an error in the Number of Sequences field, specifically:
	A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
.,.	Deleted <i>ending</i> stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patentin bug). Sequences corrected:
•	Other: corrected C1517 response
	inslited 64007 response

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95



OIPE

RAW SEQUENCE LISTING DATE: 06/27/2002 PATENT APPLICATION: US/10/073,334A TIME: 19:32:30

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF3\06272002\J073334A.raw

19 20

1 <110> APPLICANT: Steven J. Soldin
W--> 2 <120> TITLE OF INVENTION: A Novel 8.4 kDa Immunophilin
W--> 3 <130> FILE REFERENCE: 64688/155
W--> 4 <140> CURRENT APPLICATION NUMBER: 10/073,334A
C--> 5 <141> CURRENT FILING DATE: 2002-06-05
6 <150> PRIOR APPLICATION NUMBER: 09/643,723
7 <151> PRIOR FILING DATE: 2000-08-30
W--> 8 <160> NUMBER OF SEQ ID: 1
9 <170> SOFTWARE: ASCII (DOS) generic word processing
W--> 10 <210> SEQ ID NO: 1
11 <211> LENGTH: 23
12 <212> TYPE: PRT
13 <213> ORGANISM: mammal
W--> 14 <400> SEQUENCE: 1
15 Met Gln Ile Phe Val Lys Thr Leu Thr Gly

18 Lys Thr Ile Thr Leu Glu Val Glu Pro Ser

21 Asp Thr Ile

file://C:\Crf3\Outhold\VsrJ073334A.htm

VERIFICATION SUMMARY

DATE: 06/27/2002 TIME: 19:32:31

PATENT APPLICATION: US/10/073,334A

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF3\06272002\J073334A.raw

L:2 M:283 W: Missing Blank Line separator, <120> field identifier L:3 M:283 W: Missing Blank Line separator, <130> field identifier L:4 M:283 W: Missing Blank Line separator, <140> field identifier L:5 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:8 M:283 W: Missing Blank Line separator, <160> field identifier

L:10 M:283 W: Missing Blank Line separator, <210> field identifier L:14 M:283 W: Missing Blank Line separator, <400> field identifier



Does Not Comply

Corrected Diskette Needed

OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/073,334A

DATE: 06/11/2002

TIME: 16:56:27

Input Set : A:\SEQLIST.txt.6.5.02.txt
Output Set: N:\CRF3\06112002\J073334A.raw

1 <110> APPLICANT: Steven J. Soldin

W--> 2 <120> TITLE OF INVENTION: A Novel 8.4 kDa Immunophilin

W--> 3 <130> FILE REFERENCE: 64688/155

W--> 4 <140> CURRENT APPLICATION NUMBER: 10/073,334A

C--> 5 <141> CURRENT FILING DATE: 2002-06-05

6 <150> PRIOR APPLICATION NUMBER: 09/643,723

W--> 7 <151> PRIOR FILING DATE: (08/30/2000) 2000-08-50

W--> 8 <160> NUMBER OF SEQ ID: 1

9 <170> SOFTWARE: ASCII (DOS) generic word processing

## ERRORED SEQUENCES

W--> 10 <210> SEQ ID NO: 1

11 <211> LENGTH: 23

12 <212> TYPE: PRT

13 <213> ORGANISM: mammal

E--> 14 <400> SEQUENCE: 1 & MSELT

15 Met Gln Ile Phe Val Lys Thr Leu Thr Gly

6 1 . . . 5

18 Lys Thr Ile Thr Leu Glu Val Glu Pro Ser

19 15 20

21 Asp Thr Ile

**VERIFICATION SUMMARY** 

DATE: 06/11/2002

PATENT APPLICATION: US/10/073,334A

3,334A TIME: 16:56:28

Input Set : A:\SEQLIST.txt.6.5.02.txt
Output Set: N:\CRF3\06112002\J073334A.raw

L:2 M:283 W: Missing Blank Line separator, <120> field identifier L:3 M:283 W: Missing Blank Line separator, <130> field identifier L:4 M:283 W: Missing Blank Line separator, <140> field identifier

L:5 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:7 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD

L:8 M:283 W: Missing Blank Line separator, <160> field identifier L:10 M:283 W: Missing Blank Line separator, <210> field identifier

L:14 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:1 differs:0

L:14 M:283 W: Missing Blank Line separator, <400> field identifier